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**In the Specification**

Please replace the Sequence Listing in the subject application with the Sequence Listing attached hereto as **Exhibit A**.

On page 9, replace the paragraph that begins on line 6 and ends on line 16 with the following new paragraph:

-- In the present invention a method of directed evolution were applied to the lipase gene (~~Gene~~Seq ID No. ~~123~~) to isolate protein variants of the original sequence which possess increased thermostable properties. The methodology relies initially on the ability to create random variations in the original gene sequence and express the corresponding proteins in the bacteria, E.coli. The produced variants of the original sequence would have altered sequence, hence altered properties. The variants, at a proteins level , would be tested for their thermostability and those sequences which demonstrate improved thermostability would be subjected to the next round of random mutagenesis and screening. Thus by sequential accumulation of the mutants and subsequent pooling of the mutations the thermostability of lipase was improved by 200-fold at high temperature. High temperature range includes temperature ranges from 50-90 C. --

SW 9/11/07 On page <sup>10</sup>~~8~~, replace the paragraph that begins on line 28 and ends on line 4 of page <sup>11</sup>~~10~~ with the following new paragraph:

-- The naturally occurring lipase from *Bacillus lipase* has the amino acid sequence of 1-181 as given in the SEQ ID No.1 (See also SEQ ID No. 23 for corresponding gene sequence). It was discovered that the amino acid substitutions at positions 68, 71, 114, 120, 132, 144, 147 and 166 were found to be important